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SEQUENCE LISTING

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STUHMER, WALTER
BECKH, SYNNOVE
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PEREZ, ARACELI SANCHEZ
WESELOH, RUDIGER

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THEREOF

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<170> PatentIn Ver. 2.1

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 Asp Trp Asn Lys Val Ser Lys Ala Glu Ser Met Glu Thr Leu Pro Glu
 850 855 860
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 Asp Ser Gly Ile Thr Lys Ser Asp Leu Arg Leu Asp Asn Val Gly Glu
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 Ala Arg Ser Pro Gln Asp Arg Ser Pro Ile Leu Ala Glu Val Lys His
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 Ser Phe Tyr Pro Ile Pro Glu Gln Thr Leu Gln Ala Thr Val Leu Glu
 915 920 925

Val Arg His Glu Leu Lys Glu Asp Ile Lys Ala Leu Asn Ala Lys Met
 930 935 940

Thr Asn Ile Glu Lys Gln Leu Ser Glu Ile Leu Arg Ile Leu Thr Ser
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Arg Arg Ser Ser Gln Ser Pro Gln Glu Leu Phe Glu Ile Ser Arg Pro
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Gln Ser Pro Glu Ser Glu Arg Asp Ile Phe Gly Ala Ser
 980 985

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 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic DNA

<400> 5
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<210> 6
 <211> 20
 <212> DNA
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<220>
 <223> Description of Artificial Sequence: Synthetic DNA

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<210> 7
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<220>
 <223> Description of Artificial Sequence: Synthetic DNA

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<210> 8
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<220>
 <223> Description of Artificial Sequence: Synthetic DNA

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 <223> Description of Artificial Sequence: Synthetic DNA

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<220>
 <223> Description of Artificial Sequence: Synthetic DNA

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21

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21

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18

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 <213> Homo sapiens

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 <213> Homo sapiens

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<210> 15
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 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Antisense
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<400> 15
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<210> 16
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 scrambled sequence

<400> 16
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<210> 17
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 17
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<210> 18
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

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<210> 19
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 <212> DNA
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<211> 3041

<212> DNA

<213> Rattus sp.

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gtgagagtcc	cgccacgcct	gtgtccttcc	aggcagcctc	cacctccaca	gtgtcagacc	2400
acgccaaagt	gcatgcaccg	ggatctgagt	gcctagggtc	caaggcaggc	ggtggcgacc	2460
ctgccaagcg	caaaggctgg	gcccgggttca	aagatgcctg	tgggaagggt	gaggattgga	2520
acaagggtgc	caaggcagag	tccatggaga	cgcttcccga	gaggacaaaag	gcatcgggag	2580
aggccacgct	gaagaagaca	gactcctgtg	acagtggaa	caccaagagt	gacctgcgct	2640
tggacaatgt	gggtgaggcc	aggagtcccc	aggaccggag	ccccatcttg	gccgaggtca	2700
agcattcttt	ctaccccatc	cccagagcaga	cactgcaggc	cacagtgtctg	gaggtgaagc	2760
atgagctgaa	ggaagacatc	aaggccttga	atgcaaaaa	gacctccatt	gagaagcagc	2820
tgtctgagat	cctcaggata	ctcatgtcca	gagggtcctc	ccagtctccg	caggacacgt	2880
gtgaggtctc	caggccccag	tccccagagt	cagacagaga	catttttggg	gcaagctgag	2940
aggatcattt	caaaacaaac	aaacaaaaaa	atcaaagaca	aaagcctgcc	ccctgccctt	3000
gacacttcct	accgcaccaa	acacatgacc	aacaactttc	a		3041

<210> 21

<211> 960

<212> PRT

<213> Bovine sp.

<400> 21

Met	Thr	Met	Ala	Gly	Gly	Arg	Lys	Gly	Leu	Val	Ala	Pro	Gln	Asn	Thr	1	5	10	15
Phe	Leu	Glu	Asn	Ile	Val	Arg	Arg	Ser	Asn	Asp	Thr	Asn	Phe	Val	Leu	20	25	30	
Gly	Asn	Ala	Gln	Ile	Val	Asp	Trp	Pro	Ile	Val	Tyr	Ser	Asn	Asp	Gly	35	40	45	
Phe	Cys	Lys	Leu	Ser	Gly	Tyr	His	Arg	Ala	Glu	Val	Met	Gln	Lys	Ser	50	55	60	
Ser	Thr	Cys	Ser	Phe	Met	Tyr	Gly	Glu	Leu	Thr	Asp	Lys	Asp	Thr	Ile	65	70	75	80
Glu	Lys	Val	Arg	Gln	Thr	Phe	Glu	Asn	Tyr	Glu	Met	Asn	Ser	Phe	Glu	85	90	95	
Ile	Leu	Met	Tyr	Lys	Lys	Asn	Arg	Thr	Pro	Val	Trp	Phe	Phe	Val	Lys	100	105	110	
Ile	Ala	Pro	Ile	Arg	Asn	Glu	Gln	Asp	Lys	Val	Val	Leu	Phe	Leu	Cys	115	120	125	
Thr	Phe	Ser	Asp	Ile	Thr	Ala	Phe	Lys	Gln	Pro	Ile	Glu	Asp	Asp	Ser	130	135	140	
Cys	Lys	Gly	Trp	Gly	Lys	Phe	Ala	Arg	Leu	Thr	Arg	Ala	Leu	Thr	Ser	145	150	155	160
Ser	Arg	Gly	Val	Leu	Gln	Gln	Leu	Ala	Pro	Ser	Val	Gln	Lys	Gly	Glu	165	170	175	
Asn	Val	His	Lys	His	Ser	Arg	Leu	Ala	Glu	Val	Leu	Gln	Leu	Gly	Ser	180	185	190	
Asp	Ile	Leu	Pro	Gln	Tyr	Lys	Gln	Glu	Ala	Pro	Lys	Thr	Pro	Pro	His	195	200	205	
Ile	Ile	Leu	His	Tyr	Cys	Val	Phe	Lys	Thr	Thr	Trp	Asp	Trp	Ile	Ile	210	215	220	
Leu	Ile	Leu	Thr	Phe	Tyr	Thr	Ala	Ile	Leu	Val	Pro	Tyr	Asn	Val	Ser	225	230	235	240
Phe	Lys	Thr	Arg	Gln	Asn	Asn	Val	Ala	Trp	Leu	Val	Val	Asp	Ser	Ile	245	250	255	
Val	Asp	Val	Ile	Phe	Leu	Val	Asp	Ile	Val	Leu	Asn	Phe	His	Thr	Thr	260	265	270	
Phe	Val	Gly	Pro	Ala	Gly	Glu	Val	Ile	Ser	Asp	Pro	Lys	Leu	Ile	Arg	275	280	285	
Met	Asn	Tyr	Leu	Lys	Thr	Trp	Phe	Val	Ile	Asp	Leu	Leu	Ser	Cys	Leu	290	295	300	

Pro	Tyr	Asp	Val	Ile	Asn	Ala	Phe	Glu	Asn	Val	Asp	Glu	Gly	Ile	Ser	305	310	315	320
Ser	Leu	Phe	Ser	Ser	Leu	Lys	Val	Val	Arg	Leu	Leu	Arg	Leu	Gly	Arg	325	330	335	
Val	Ala	Arg	Lys	Leu	Asp	His	Tyr	Ile	Glu	Tyr	Gly	Ala	Ala	Val	Leu	340	345	350	
Val	Leu	Leu	Val	Cys	Val	Phe	Gly	Leu	Ala	Ala	His	Trp	Met	Ala	Cys	355	360	365	
Ile	Trp	Tyr	Ser	Ile	Gly	Asp	Tyr	Glu	Ile	Phe	Asp	Glu	Asp	Thr	Lys	370	375	380	
Thr	Ile	Arg	Asn	Asn	Ser	Trp	Leu	Tyr	Gln	Leu	Ala	Met	Asp	Ile	Gly	385	390	395	400
Thr	Pro	Tyr	Gln	Phe	Asn	Gly	Ser	Gly	Ser	Gly	Lys	Trp	Glu	Gly	Gly	405	410	415	
Pro	Ser	Lys	Asn	Ser	Val	Tyr	Ile	Ser	Ser	Leu	Tyr	Phe	Thr	Met	Thr	420	425	430	
Ser	Leu	Thr	Ser	Val	Gly	Phe	Gly	Asn	Ile	Ala	Pro	Ser	Thr	Asp	Ile	435	440	445	
Glu	Lys	Ile	Phe	Ala	Val	Ala	Ile	Met	Met	Ile	Gly	Ser	Leu	Leu	Tyr	450	455	460	
Ala	Thr	Ile	Phe	Gly	Asn	Val	Thr	Thr	Ile	Phe	Gln	Gln	Met	Tyr	Ala	465	470	475	480
Asn	Thr	Asn	Arg	Tyr	His	Glu	Met	Leu	Asn	Ser	Val	Arg	Asp	Phe	Leu	485	490	495	
Lys	Leu	Tyr	Gln	Val	Pro	Lys	Gly	Leu	Ser	Glu	Arg	Val	Met	Asp	Tyr	500	505	510	
Ile	Val	Ser	Thr	Trp	Ser	Met	Ser	Arg	Gly	Ile	Asp	Thr	Glu	Lys	Val	515	520	525	
Leu	Gln	Ile	Cys	Pro	Lys	Asp	Met	Arg	Ala	Asp	Ile	Cys	Val	His	Leu	530	535	540	
Asn	Arg	Lys	Val	Phe	Lys	Glu	His	Pro	Ala	Phe	Arg	Leu	Ala	Ser	Asp	545	550	555	560
Gly	Cys	Leu	Arg	Ala	Leu	Ala	Met	Glu	Phe	Gln	Thr	Val	His	Cys	Ala	565	570	575	
Pro	Gly	Asp	Leu	Ile	Tyr	His	Ala	Gly	Glu	Ser	Val	Asp	Ser	Leu	Cys	580	585	590	
Phe	Val	Val	Ser	Gly	Ser	Leu	Glu	Val	Ile	Gln	Asp	Asp	Glu	Val	Val	595	600	605	

Ala	Ile	Leu	Gly	Lys	Gly	Asp	Val	Phe	Gly	Asp	Val	Phe	Trp	Lys	Glu	610	615	620
Ala	Thr	Leu	Ala	Gln	Ser	Cys	Ala	Asn	Val	Arg	Ala	Leu	Thr	Tyr	Cys	625	630	635
Asp	Leu	His	Val	Ile	Lys	Arg	Asp	Ala	Leu	Gln	Lys	Val	Leu	Glu	Phe	645	650	655
Tyr	Thr	Ala	Phe	Ser	His	Ser	Phe	Ser	Arg	Asn	Leu	Ile	Leu	Thr	Tyr	660	665	670
Asn	Leu	Arg	Lys	Arg	Ile	Val	Phe	Arg	Lys	Ile	Ser	Asp	Val	Lys	Arg	675	680	685
Glu	Glu	Glu	Glu	Arg	Met	Lys	Arg	Lys	Asn	Glu	Ala	Pro	Leu	Ile	Leu	690	695	700
Pro	Pro	Asp	His	Pro	Val	Arg	Arg	Leu	Phe	Gln	Arg	Phe	Arg	Gln	Gln	705	710	715
Lys	Glu	Ala	Arg	Leu	Ala	Ala	Glu	Arg	Gly	Gly	Arg	Asp	Leu	Asp	Asp	725	730	735
Leu	Asp	Val	Glu	Lys	Gly	Ser	Val	Leu	Thr	Glu	His	Ser	His	His	Gly	740	745	750
Leu	Ala	Lys	Ala	Ser	Val	Val	Thr	Val	Arg	Glu	Ser	Pro	Ala	Thr	Pro	755	760	765
Val	Ala	Phe	Pro	Ala	Ala	Ala	Ala	Pro	Ala	Gly	Leu	Asp	His	Ala	Arg	770	775	780
Leu	Gln	Ala	Pro	Gly	Ala	Glu	Gly	Leu	Gly	Pro	Lys	Ala	Gly	Gly	Ala	785	790	795
Asp	Cys	Ala	Lys	Arg	Lys	Gly	Trp	Ala	Arg	Phe	Lys	Asp	Ala	Cys	Gly	805	810	815
Gln	Ala	Glu	Asp	Trp	Ser	Lys	Val	Ser	Lys	Ala	Glu	Ser	Met	Glu	Thr	820	825	830
Leu	Pro	Glu	Arg	Thr	Lys	Ala	Ala	Gly	Glu	Ala	Thr	Leu	Lys	Lys	Thr	835	840	845
Asp	Ser	Cys	Asp	Ser	Gly	Ile	Thr	Lys	Ser	Asp	Leu	Arg	Leu	Asp	Asn	850	855	860
Val	Gly	Glu	Ala	Arg	Ser	Pro	Gln	Asp	Arg	Ser	Pro	Ile	Leu	Ala	Glu	865	870	875
Val	Lys	His	Ser	Phe	Tyr	Pro	Ile	Pro	Glu	Gln	Thr	Leu	Gln	Ala	Ala	885	890	895
Val	Leu	Glu	Val	Lys	His	Glu	Leu	Lys	Glu	Asp	Ile	Lys	Ala	Leu	Ser	900	905	910

Thr Lys Met Thr Ser Ile Glu Lys Gln Leu Ser Glu Ile Leu Arg Ile
 915 920 925
 Leu Thr Ser Arg Arg Ser Ser Gln Ser Pro Gln Glu Leu Phe Glu Ile
 930 935 940
 Ser Arg Pro Gln Ser Pro Glu Ser Glu Arg Asp Ile Phe Gly Ala Ser
 945 950 955 960

<210> 22
 <211> 987
 <212> PRT
 <213> Bovine sp.

<400> 22
 Met Thr Met Ala Gly Gly Arg Lys Gly Leu Val Ala Pro Gln Asn Thr
 1 5 10 15
 Phe Leu Glu Asn Ile Val Arg Arg Ser Asn Asp Thr Asn Phe Val Leu
 20 25 30
 Gly Asn Ala Gln Ile Val Asp Trp Pro Ile Val Tyr Ser Asn Asp Gly
 35 40 45
 Phe Cys Lys Leu Ser Gly Tyr His Arg Ala Glu Val Met Gln Lys Ser
 50 55 60
 Ser Thr Cys Ser Phe Met Tyr Gly Glu Leu Thr Asp Lys Asp Thr Ile
 65 70 75 80
 Glu Lys Val Arg Gln Thr Phe Glu Asn Tyr Glu Met Asn Ser Phe Glu
 85 90 95
 Ile Leu Met Tyr Lys Lys Asn Arg Thr Pro Val Trp Phe Phe Val Lys
 100 105 110
 Ile Ala Pro Ile Arg Asn Glu Gln Asp Lys Val Val Leu Phe Leu Cys
 115 120 125
 Thr Phe Ser Asp Ile Thr Ala Phe Lys Gln Pro Ile Glu Asp Asp Ser
 130 135 140
 Cys Lys Gly Trp Gly Lys Phe Ala Arg Leu Thr Arg Ala Leu Thr Ser
 145 150 155 160
 Ser Arg Gly Val Leu Gln Gln Leu Ala Pro Ser Val Gln Lys Gly Glu
 165 170 175
 Asn Val His Lys His Ser Arg Leu Ala Glu Val Leu Gln Leu Gly Ser
 180 185 190
 Asp Ile Leu Pro Gln Tyr Lys Gln Glu Ala Pro Lys Thr Pro Pro His
 195 200 205
 Ile Ile Leu His Tyr Cys Val Phe Lys Thr Thr Trp Asp Trp Ile Ile
 210 215 220

Leu	Ile	Leu	Thr	Phe	Tyr	Thr	Ala	Ile	Leu	Val	Pro	Tyr	Asn	Val	Ser
225					230					235					240
Phe	Lys	Thr	Arg	Gln	Asn	Asn	Val	Ala	Trp	Leu	Val	Val	Asp	Ser	Ile
				245					250					255	
Val	Asp	Val	Ile	Phe	Leu	Val	Asp	Ile	Val	Leu	Asn	Phe	His	Thr	Thr
			260					265					270		
Phe	Val	Gly	Pro	Ala	Gly	Glu	Val	Ile	Ser	Asp	Pro	Lys	Leu	Ile	Arg
		275					280					285			
Met	Asn	Tyr	Leu	Lys	Thr	Trp	Phe	Val	Ile	Asp	Leu	Leu	Ser	Cys	Leu
	290					295					300				
Pro	Tyr	Asp	Val	Ile	Asn	Ala	Phe	Glu	Asn	Val	Asp	Glu	Val	Ser	Ala
305					310					315					320
Phe	Met	Gly	Asp	Pro	Gly	Lys	Ile	Gly	Phe	Ala	Asp	Gln	Ile	Pro	Pro
				325					330					335	
Pro	Leu	Glu	Gly	Arg	Glu	Ser	Gln	Gly	Ile	Ser	Ser	Leu	Phe	Ser	Ser
			340					345					350		
Leu	Lys	Val	Val	Arg	Leu	Leu	Arg	Leu	Gly	Arg	Val	Ala	Arg	Lys	Leu
		355					360					365			
Asp	His	Tyr	Ile	Glu	Tyr	Gly	Ala	Ala	Val	Leu	Val	Leu	Leu	Val	Cys
	370					375					380				
Val	Phe	Gly	Leu	Ala	Ala	His	Trp	Met	Ala	Cys	Ile	Trp	Tyr	Ser	Ile
385					390					395					400
Gly	Asp	Tyr	Glu	Ile	Phe	Asp	Glu	Asp	Thr	Lys	Thr	Ile	Arg	Asn	Asn
				405					410					415	
Ser	Trp	Leu	Tyr	Gln	Leu	Ala	Met	Asp	Ile	Gly	Thr	Pro	Tyr	Gln	Phe
			420					425					430		
Asn	Gly	Ser	Gly	Ser	Gly	Lys	Trp	Glu	Gly	Gly	Pro	Ser	Lys	Asn	Ser
		435					440					445			
Val	Tyr	Ile	Ser	Ser	Leu	Tyr	Phe	Thr	Met	Thr	Ser	Leu	Thr	Ser	Val
	450					455					460				
Gly	Phe	Gly	Asn	Ile	Ala	Pro	Ser	Thr	Asp	Ile	Glu	Lys	Ile	Phe	Ala
465					470					475					480
Val	Ala	Ile	Met	Met	Ile	Gly	Ser	Leu	Leu	Tyr	Ala	Thr	Ile	Phe	Gly
				485					490					495	
Asn	Val	Thr	Thr	Ile	Phe	Gln	Gln	Met	Tyr	Ala	Asn	Thr	Asn	Arg	Tyr
			500					505					510		
His	Glu	Met	Leu	Asn	Ser	Val	Arg	Asp	Phe	Leu	Lys	Leu	Tyr	Gln	Val
		515					520					525			

Pro Lys Gly Leu Ser Glu Arg Val Met Asp Tyr Ile Val Ser Thr Trp
 530 535 540
 Ser Met Ser Arg Gly Ile Asp Thr Glu Lys Val Leu Gln Ile Cys Pro
 545 550 555 560
 Lys Asp Met Arg Ala Asp Ile Cys Val His Leu Asn Arg Lys Val Phe
 565 570 575
 Lys Glu His Pro Ala Phe Arg Leu Ala Ser Asp Gly Cys Leu Arg Ala
 580 585 590
 Leu Ala Met Glu Phe Gln Thr Val His Cys Ala Pro Gly Asp Leu Ile
 595 600 605
 Tyr His Ala Gly Glu Ser Val Asp Ser Leu Cys Phe Val Val Ser Gly
 610 615 620
 Ser Leu Glu Val Ile Gln Asp Asp Glu Val Val Ala Ile Leu Gly Lys
 625 630 635 640
 Gly Asp Val Phe Gly Asp Val Phe Trp Lys Glu Ala Thr Leu Ala Gln
 645 650 655
 Ser Cys Ala Asn Val Arg Ala Leu Thr Tyr Cys Asp Leu His Val Ile
 660 665 670
 Lys Arg Asp Ala Leu Gln Lys Val Leu Glu Phe Tyr Thr Ala Phe Ser
 675 680 685
 His Ser Phe Ser Arg Asn Leu Ile Leu Thr Tyr Asn Leu Arg Lys Arg
 690 695 700
 Ile Val Phe Arg Lys Ile Ser Asp Val Lys Arg Glu Glu Glu Glu Arg
 705 710 715 720
 Met Lys Arg Lys Asn Glu Ala Pro Leu Ile Leu Pro Pro Asp His Pro
 725 730 735
 Val Arg Arg Leu Phe Gln Arg Phe Arg Gln Gln Lys Glu Ala Arg Leu
 740 745 750
 Ala Ala Glu Arg Gly Gly Arg Asp Leu Asp Asp Leu Asp Val Glu Lys
 755 760 765
 Gly Ser Val Leu Thr Glu His Ser His His Gly Leu Ala Lys Ala Ser
 770 775 780
 Val Val Thr Val Arg Glu Ser Pro Ala Thr Pro Val Ala Phe Pro Ala
 785 790 795 800
 Ala Ala Ala Pro Ala Gly Leu Asp His Ala Arg Leu Gln Ala Pro Gly
 805 810 815
 Ala Glu Gly Leu Gly Pro Lys Ala Gly Gly Ala Asp Cys Ala Lys Arg
 820 825 830

Lys Gly Trp Ala Arg Phe Lys Asp Ala Cys Gly Gln Ala Glu Asp Trp
 835 840 845
 Ser Lys Val Ser Lys Ala Glu Ser Met Glu Thr Leu Pro Glu Arg Thr
 850 855 860
 Lys Ala Ala Gly Glu Ala Thr Leu Lys Lys Thr Asp Ser Cys Asp Ser
 865 870 875 880
 Gly Ile Thr Lys Ser Asp Leu Arg Leu Asp Asn Val Gly Glu Ala Arg
 885 890 895
 Ser Pro Gln Asp Arg Ser Pro Ile Leu Ala Glu Val Lys His Ser Phe
 900 905 910
 Tyr Pro Ile Pro Glu Gln Thr Leu Gln Ala Ala Val Leu Glu Val Lys
 915 920 925
 His Glu Leu Lys Glu Asp Ile Lys Ala Leu Ser Thr Lys Met Thr Ser
 930 935 940
 Ile Glu Lys Gln Leu Ser Glu Ile Leu Arg Ile Leu Thr Ser Arg Arg
 945 950 955 960
 Ser Ser Gln Ser Pro Gln Glu Leu Phe Glu Ile Ser Arg Pro Gln Ser
 965 970 975
 Pro Glu Ser Glu Arg Asp Ile Phe Gly Ala Ser
 980 985

<210> 23
 <211> 989
 <212> PRT
 <213> Mus sp.

<400> 23

Met Thr Met Ala Gly Gly Arg Lys Gly Leu Val Ala Pro Gln Asn Thr
 1 5 10 15
 Phe Leu Glu Asn Ile Val Arg Arg Ser Asn Asp Thr Asn Phe Val Leu
 20 25 30
 Gly Asn Ala Gln Ile Val Asp Trp Pro Ile Val Tyr Ser Asn Asp Gly
 35 40 45
 Phe Cys Lys Leu Ser Gly Tyr His Arg Ala Glu Val Met Gln Lys Ser
 50 55 60
 Ser Ala Cys Ser Phe Met Tyr Gly Glu Leu Thr Asp Lys Asp Thr Val
 65 70 75 80
 Glu Lys Val Arg Gln Thr Phe Glu Asn Tyr Glu Met Asn Ser Phe Glu
 85 90 95
 Ile Leu Met Tyr Lys Lys Asn Arg Thr Pro Val Trp Phe Phe Val Lys
 100 105 110

Ile	Ala	Pro	Ile	Arg	Asn	Glu	Gln	Asp	Lys	Val	Val	Leu	Phe	Leu	Cys
		115					120					125			
Thr	Phe	Ser	Asp	Ile	Thr	Ala	Phe	Lys	Gln	Pro	Ile	Glu	Asp	Asp	Ser
	130					135					140				
Cys	Lys	Gly	Trp	Gly	Lys	Phe	Ala	Arg	Leu	Thr	Arg	Ala	Leu	Thr	Ser
145					150					155					160
Ser	Arg	Gly	Val	Leu	Gln	Gln	Leu	Ala	Pro	Ser	Val	Gln	Lys	Gly	Glu
			165						170					175	
Asn	Val	His	Lys	His	Ser	Arg	Leu	Ala	Glu	Val	Leu	Gln	Leu	Gly	Ser
		180						185					190		
Asp	Ile	Leu	Pro	Gln	Tyr	Lys	Gln	Glu	Ala	Pro	Lys	Thr	Pro	Pro	His
		195					200					205			
Ile	Ile	Leu	His	Tyr	Cys	Val	Phe	Lys	Thr	Thr	Trp	Asp	Trp	Ile	Ile
	210					215					220				
Leu	Ile	Leu	Thr	Phe	Tyr	Thr	Ala	Ile	Leu	Val	Pro	Tyr	Asn	Val	Ser
225					230					235					240
Phe	Lys	Thr	Arg	Gln	Asn	Asn	Val	Ala	Trp	Leu	Val	Val	Asp	Ser	Ile
				245					250					255	
Val	Asp	Val	Ile	Phe	Leu	Val	Asp	Ile	Val	Leu	Asn	Phe	His	Thr	Thr
		260						265					270		
Phe	Val	Gly	Pro	Ala	Gly	Glu	Val	Ile	Ser	Asp	Pro	Lys	Leu	Ile	Arg
		275					280					285			
Met	Asn	Tyr	Leu	Lys	Thr	Trp	Phe	Val	Ile	Asp	Leu	Leu	Ser	Cys	Leu
	290					295					300				
Pro	Tyr	Asp	Val	Ile	Asn	Ala	Phe	Glu	Asn	Val	Asp	Glu	Val	Ser	Ala
305					310					315					320
Phe	Met	Gly	Asp	Pro	Gly	Lys	Ile	Gly	Phe	Ala	Asp	Gln	Ile	Pro	Pro
				325					330					335	
Pro	Leu	Glu	Gly	Arg	Glu	Ser	Gln	Gly	Ile	Ser	Ser	Leu	Phe	Ser	Ser
			340					345					350		
Leu	Lys	Val	Val	Arg	Leu	Leu	Arg	Leu	Gly	Arg	Val	Ala	Arg	Lys	Leu
		355					360					365			
Asp	His	Tyr	Ile	Glu	Tyr	Gly	Ala	Ala	Val	Leu	Val	Leu	Leu	Val	Cys
	370					375					380				
Val	Phe	Gly	Leu	Ala	Ala	His	Trp	Met	Ala	Cys	Ile	Trp	Tyr	Ser	Ile
385					390					395					400
Gly	Asp	Tyr	Glu	Ile	Phe	Asp	Glu	Asp	Thr	Lys	Thr	Ile	Arg	Asn	Asn
				405					410					415	

Ser Trp Leu Tyr Gln Leu Ala Leu Asp Ile Gly Thr Pro Tyr Gln Phe
 420 425 430
 Asn Gly Ser Gly Ser Gly Lys Trp Glu Gly Gly Pro Ser Lys Asn Ser
 435 440 445
 Val Tyr Ile Ser Ser Leu Tyr Phe Thr Met Thr Ser Leu Thr Ser Val
 450 455 460
 Gly Phe Gly Asn Ile Ala Pro Ser Thr Asp Ile Glu Lys Ile Phe Ala
 465 470 475 480
 Val Ala Ile Met Met Ile Gly Ser Leu Leu Tyr Ala Thr Ile Phe Gly
 485 490 495
 Asn Val Thr Thr Ile Phe Gln Gln Met Tyr Ala Asn Thr Asn Arg Tyr
 500 505 510
 His Glu Met Leu Asn Ser Val Arg Asp Phe Leu Lys Leu Tyr Gln Val
 515 520 525
 Pro Lys Gly Leu Ser Glu Arg Val Met Asp Tyr Ile Val Ser Thr Trp
 530 535 540
 Ser Met Ser Arg Gly Ile Asp Thr Glu Lys Val Leu Gln Ile Cys Pro
 545 550 555 560
 Lys Asp Met Arg Ala Asp Ile Cys Val His Leu Asn Arg Lys Val Phe
 565 570 575
 Lys Glu His Pro Ala Phe Arg Leu Ala Ser Asp Gly Cys Leu Arg Ala
 580 585 590
 Leu Ala Met Glu Phe Gln Thr Val His Cys Ala Pro Gly Asp Leu Ile
 595 600 605
 Tyr His Ala Gly Glu Ser Val Asp Ser Leu Cys Phe Val Val Ser Gly
 610 615 620
 Ser Leu Glu Val Ile Gln Asp Asp Glu Val Val Ala Ile Leu Gly Lys
 625 630 635 640
 Gly Asp Val Phe Gly Asp Val Phe Trp Lys Glu Ala Thr Leu Ala Gln
 645 650 655
 Ser Cys Ala Asn Val Arg Ala Leu Thr Tyr Cys Asp Leu His Val Ile
 660 665 670
 Lys Arg Asp Ala Leu Gln Lys Val Leu Glu Phe Tyr Thr Ala Phe Ser
 675 680 685
 His Ser Phe Ser Arg Asn Leu Ile Leu Thr Tyr Asn Leu Arg Lys Arg
 690 695 700
 Ile Val Phe Arg Lys Ile Ser Asp Val Lys Arg Glu Glu Glu Glu Arg
 705 710 715 720

Met Lys Arg Lys Asn Glu Ala Pro Leu Ile Leu Pro Pro Asp His Pro
 725 730 735
 Val Arg Arg Leu Phe Gln Arg Phe Arg Gln Gln Lys Glu Ala Arg Leu
 740 745 750
 Ala Ala Glu Arg Gly Gly Arg Asp Leu Asp Asp Leu Asp Val Glu Lys
 755 760 765
 Gly Asn Ala Leu Thr Asp His Thr Ser Ala Asn His Gly Leu Ala Lys
 770 775 780
 Ala Ser Val Val Thr Val Arg Glu Ser Pro Ala Thr Pro Val Ala Phe
 785 790 795 800
 Gln Ala Ala Thr Thr Ser Thr Met Ser Asp His Ala Lys Leu His Ala
 805 810 815
 Pro Gly Ser Glu Cys Leu Gly Pro Lys Ala Val Ser Cys Asp Pro Ala
 820 825 830
 Lys Arg Lys Gly Trp Ala Arg Phe Lys Asp Ala Cys Gly Gln Ala Glu
 835 840 845
 Asp Trp Ser Lys Val Ser Lys Ala Glu Ser Met Glu Thr Leu Pro Glu
 850 855 860
 Arg Thr Lys Ala Pro Gly Glu Ala Thr Leu Lys Lys Thr Asp Ser Cys
 865 870 875 880
 Asp Ser Gly Ile Thr Lys Ser Asp Leu Arg Leu Asp Asn Val Gly Glu
 885 890 895
 Thr Arg Ser Pro Gln Asp Arg Ser Pro Ile Leu Ala Glu Val Lys His
 900 905 910
 Ser Phe Tyr Pro Ile Pro Glu Gln Thr Leu Gln Ala Ala Val Leu Glu
 915 920 925
 Val Lys Tyr Glu Leu Lys Glu Asp Ile Lys Ala Leu Asn Ala Lys Met
 930 935 940
 Thr Ser Ile Glu Lys Gln Leu Ser Glu Ile Leu Arg Ile Leu Met Ser
 945 950 955 960
 Arg Gly Ser Ala Gln Ser Pro Gln Glu Thr Gly Glu Ile Ser Arg Pro
 965 970 975
 Gln Ser Pro Glu Ser Asp Arg Asp Ile Phe Gly Ala Ser
 980 985

<210> 24

<211> 962

<212> PRT

<213> Rattus sp.

<400> 24

Met	Thr	Met	Ala	Gly	Gly	Arg	Lys	Gly	Leu	Val	Ala	Pro	Gln	Asn	Thr	1	5	10	15
Phe	Leu	Glu	Asn	Ile	Val	Arg	Arg	Ser	Asn	Asp	Thr	Asn	Phe	Val	Leu	20	25	30	
Gly	Asn	Ala	Gln	Ile	Val	Asp	Trp	Pro	Ile	Val	Tyr	Ser	Asn	Asp	Gly	35	40	45	
Phe	Cys	Lys	Leu	Ser	Gly	Tyr	His	Arg	Ala	Glu	Val	Met	Gln	Lys	Ser	50	55	60	
Ser	Ala	Cys	Ser	Phe	Met	Tyr	Gly	Glu	Leu	Thr	Asp	Lys	Asp	Thr	Val	65	70	75	80
Glu	Lys	Val	Arg	Gln	Thr	Phe	Glu	Asn	Tyr	Glu	Met	Asn	Ser	Phe	Glu	85	90	95	
Ile	Leu	Met	Tyr	Lys	Lys	Asn	Arg	Thr	Pro	Val	Trp	Phe	Phe	Val	Lys	100	105	110	
Ile	Ala	Pro	Ile	Arg	Asn	Glu	Gln	Asp	Lys	Val	Val	Leu	Phe	Leu	Cys	115	120	125	
Thr	Phe	Ser	Asp	Ile	Thr	Ala	Phe	Lys	Gln	Pro	Ile	Glu	Asp	Asp	Ser	130	135	140	
Cys	Lys	Gly	Trp	Gly	Lys	Phe	Ala	Arg	Leu	Thr	Arg	Ala	Leu	Thr	Ser	145	150	155	160
Ser	Arg	Gly	Val	Leu	Gln	Gln	Leu	Ala	Pro	Ser	Val	Gln	Lys	Gly	Glu	165	170	175	
Asn	Val	His	Lys	His	Ser	Arg	Leu	Ala	Glu	Val	Leu	Gln	Leu	Gly	Ser	180	185	190	
Asp	Ile	Leu	Pro	Gln	Tyr	Lys	Gln	Glu	Ala	Pro	Lys	Thr	Pro	Pro	His	195	200	205	
Ile	Ile	Leu	His	Tyr	Cys	Val	Phe	Lys	Thr	Thr	Trp	Asp	Trp	Ile	Ile	210	215	220	
Leu	Ile	Leu	Thr	Phe	Tyr	Thr	Ala	Ile	Leu	Val	Pro	Tyr	Asn	Val	Ser	225	230	235	240
Phe	Lys	Thr	Arg	Gln	Asn	Asn	Val	Ala	Trp	Leu	Val	Val	Asp	Ser	Ile	245	250	255	
Val	Asp	Val	Ile	Phe	Leu	Val	Asp	Ile	Val	Leu	Asn	Phe	His	Thr	Thr	260	265	270	
Phe	Val	Gly	Pro	Ala	Gly	Glu	Val	Ile	Ser	Asp	Pro	Lys	Leu	Ile	Arg	275	280	285	
Met	Asn	Tyr	Leu	Lys	Thr	Trp	Phe	Val	Ile	Asp	Leu	Leu	Ser	Cys	Leu	290	295	300	

Pro Tyr Asp Val Ile Asn Ala Phe Glu Asn Val Asp Glu Gly Ile Ser
 305 310 315 320
 Ser Leu Phe Ser Ser Leu Lys Val Val Arg Leu Leu Arg Leu Gly Arg
 325 330 335
 Val Ala Arg Lys Leu Asp His Tyr Ile Glu Tyr Gly Ala Ala Val Leu
 340 345 350
 Val Leu Leu Val Cys Val Phe Gly Leu Ala Ala His Trp Met Ala Cys
 355 360 365
 Ile Trp Tyr Ser Ile Gly Asp Tyr Glu Ile Phe Asp Glu Asp Thr Lys
 370 375 380
 Thr Ile Arg Asn Asn Ser Trp Leu Tyr Gln Leu Ala Leu Asp Ile Gly
 385 390 395 400
 Thr Pro Tyr Gln Phe Asn Gly Ser Gly Ser Gly Lys Trp Glu Gly Gly
 405 410 415
 Pro Ser Lys Asn Ser Val Tyr Ile Ser Ser Leu Tyr Phe Thr Met Thr
 420 425 430
 Ser Leu Thr Ser Val Gly Phe Gly Asn Ile Ala Pro Ser Thr Asp Ile
 435 440 445
 Glu Lys Ile Phe Ala Val Ala Ile Met Met Ile Gly Ser Leu Leu Tyr
 450 455 460
 Ala Thr Ile Phe Gly Asn Val Thr Thr Ile Phe Gln Gln Met Tyr Ala
 465 470 475 480
 Asn Thr Asn Arg Tyr His Glu Met Leu Asn Ser Val Arg Asp Phe Leu
 485 490 495
 Lys Leu Tyr Gln Val Pro Lys Gly Leu Ser Glu Arg Val Met Asp Tyr
 500 505 510
 Ile Val Ser Thr Trp Ser Met Ser Arg Gly Ile Asp Thr Glu Lys Val
 515 520 525
 Leu Gln Ile Cys Pro Lys Asp Met Arg Ala Asp Ile Cys Val His Leu
 530 535 540
 Asn Arg Lys Val Phe Lys Glu His Pro Ala Phe Arg Leu Ala Ser Asp
 545 550 555 560
 Gly Cys Leu Arg Ala Leu Ala Met Glu Phe Gln Thr Val His Cys Ala
 565 570 575
 Pro Gly Asp Leu Ile Tyr His Ala Gly Glu Ser Val Asp Ser Leu Cys
 580 585 590
 Phe Val Val Ser Gly Ser Leu Glu Val Ile Gln Asp Asp Glu Val Val
 595 600 605

Ala	Ile	Leu	Gly	Lys	Gly	Asp	Val	Phe	Gly	Asp	Val	Phe	Trp	Lys	Glu
610						615					620				
Ala	Thr	Leu	Ala	Gln	Ser	Cys	Ala	Asn	Val	Arg	Ala	Leu	Thr	Tyr	Cys
625					630					635					640
Asp	Leu	His	Val	Ile	Lys	Arg	Asp	Ala	Leu	Gln	Lys	Val	Leu	Glu	Phe
				645					650					655	
Tyr	Thr	Ala	Phe	Ser	His	Ser	Phe	Ser	Arg	Asn	Leu	Ile	Leu	Thr	Tyr
			660					665					670		
Asn	Leu	Arg	Lys	Arg	Ile	Val	Phe	Arg	Lys	Ile	Ser	Asp	Val	Lys	Arg
		675					680					685			
Glu	Glu	Glu	Glu	Arg	Met	Lys	Arg	Lys	Asn	Glu	Ala	Pro	Leu	Ile	Leu
	690					695					700				
Pro	Pro	Asp	His	Pro	Val	Arg	Arg	Leu	Phe	Gln	Arg	Phe	Arg	Gln	Gln
705					710					715					720
Lys	Glu	Ala	Arg	Leu	Ala	Ala	Glu	Arg	Gly	Gly	Arg	Asp	Leu	Asp	Asp
				725					730					735	
Leu	Asp	Val	Glu	Lys	Gly	Asn	Ala	Leu	Thr	Asp	His	Thr	Ser	Ala	Asn
			740					745					750		
His	Gly	Leu	Ala	Lys	Ala	Ser	Val	Val	Thr	Val	Arg	Glu	Ser	Pro	Ala
		755					760					765			
Thr	Pro	Val	Ala	Phe	Gln	Ala	Ala	Ser	Thr	Ser	Thr	Val	Ser	Asp	His
	770					775					780				
Ala	Lys	Leu	His	Ala	Pro	Gly	Ser	Glu	Cys	Leu	Gly	Pro	Lys	Ala	Gly
785					790				795						800
Gly	Gly	Asp	Pro	Ala	Lys	Arg	Lys	Gly	Trp	Ala	Arg	Phe	Lys	Asp	Ala
				805					810					815	
Cys	Gly	Gln	Ala	Glu	Asp	Trp	Ser	Lys	Val	Ser	Lys	Ala	Glu	Ser	Met
			820					825					830		
Glu	Thr	Leu	Pro	Glu	Arg	Thr	Lys	Ala	Ala	Gly	Glu	Ala	Thr	Leu	Lys
		835					840					845			
Lys	Thr	Asp	Ser	Cys	Asp	Ser	Gly	Ile	Thr	Lys	Ser	Asp	Leu	Arg	Leu
	850					855					860				
Asp	Asn	Val	Gly	Glu	Ala	Arg	Ser	Pro	Gln	Asp	Arg	Ser	Pro	Ile	Leu
865					870					875					880
Ala	Glu	Val	Lys	His	Ser	Phe	Tyr	Pro	Ile	Pro	Glu	Gln	Thr	Leu	Gln
				885					890					895	
Ala	Thr	Val	Leu	Glu	Val	Lys	Tyr	Glu	Leu	Lys	Glu	Asp	Ile	Lys	Ala
			900					905					910		

Leu Asn Ala Lys Met Thr Ser Ile Glu Lys Gln Leu Ser Glu Ile Leu
915 920 925

Arg Ile Leu Met Ser Arg Gly Ser Ser Gln Ser Pro Gln Asp Thr Cys
930 935 940

Glu Val Ser Arg Pro Gln Ser Pro Glu Ser Asp Arg Asp Ile Phe Gly
945 950 955 960

Ala Ser

C15
CONT